

SEQ SEARCH SEQ ID NO:2 (STIC):

```
us-09-701-586d-2.rag nothing 1/20/2010 rh
us-09-701-586d-2.rai nothing 1/20/2010 rh
us-09-701-586d-2.rapb nothing 1/20/2010 rh
us-09-701-586d-2.rpr nothing 1/20/2010 rh
us-09-701-586d-2.rsp nothing 1/20/2010 rh
us-09-701-586d-2.rapm nothing 1/20/2010 rh
us-09-701-586d-2.rapn nothing 1/20/2010 rh
us-09-701-586d-2.rspt nothing 1/20/2010 rh
```

RESULT 8

```

US-10-369-378-2
; Sequence 2, Application US/10369378
; GENERAL INFORMATION:
; APPLICANT: Christenson, Erik
; APPLICANT: DeMaggio, Anthony J
; APPLICANT: Goldman, Phyllis S
; APPLICANT: McElligott, David L
; TITLE OF INVENTION: Human Poly(ADP-Ribose) Polymerase 2 Materials and
; TITLE OF INVENTION: Methods
; FILE REFERENCE: 27866/36544
; CURRENT APPLICATION NUMBER: US/10/369,378
; CURRENT FILING DATE: 2003-02-19
; PRIOR APPLICATION NUMBER: US/09/596,248D
; PRIOR FILING DATE: 2000-06-16
; PRIOR APPLICATION NUMBER: 60/139,543
; PRIOR FILING DATE: 1999-06-16
; NUMBER OF SEQ ID NOS: 68
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 583
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-369-378-2

```

Query Match 99.4%; Score 2981.5; DB 29; Length 583;
Best Local Similarity 97.8%; Pred. No. 3.3e-262;
Matches 570; Conservative 0; Mismatches 0; Indels 13; Gaps 1;

Qy	1	MAARRRRSTGGGRARALNESKRVNNGNTAPEDSSPAKKTRRCQRQESKKMPVAGGKANKD	60
Db	1	MAARRRRSTGGGRARALNESKRVNNGNTAPEDSSPAKKTRRCQRQESKKMPVAGGKANKD	60
Qy	61	RTEDKQD-----ESVKALLLKGPVDPECTAKVGKAHVYCEGNDVYDVMLN	107
Db	61	RTEDKQDGMPPGRSWASKRVSESVKALLLKGPVDPECTAKVGKAHVYCEGNDVYDVMLN	120

Qy	108	QTNLQFNNNKYYLIQLLEDDAQRNFSVWMRWGRVGMKGQHSVLVACSGNLNKAKEIFQKKF	167
Db	121	QTNLQFNNNKYYLIQLLEDDAQRNFSVWMRWGRVGMKGQHSVLVACSGNLNKAKEIFQKKF	180
Qy	168	LDKTKNNWEDREKFVKPGKYDMLQMDYATNTQDEEETKKEESLKSPLKPESQLDLRVQE	227
Db	181	LDKTKNNWEDREKFVKPGKYDMLQMDYATNTQDEEETKKEESLKSPLKPESQLDLRVQE	240
Qy	228	LIKLICNVQAMEEMMEMKYNTKKAPLGKLTVAQIKAGYQSLKKIEDCIRAGQHGRALME	287
Db	241	LIKLICNVQAMEEMMEMKYNTKKAPLGKLTVAQIKAGYQSLKKIEDCIRAGQHGRALME	300
Qy	288	ACNEFYTRIPHDFGLRTPPLIRTQKELSEKIQLLEALGDIEIAIKLVKTELQSPEHPLDQ	347
Db	301	ACNEFYTRIPHDFGLRTPPLIRTQKELSEKIQLLEALGDIEIAIKLVKTELQSPEHPLDQ	360
Qy	348	HYRNLHCALRPLDHESYEFKVISQYLQSTHAPTHSDYTMLLDLFEVEKDGEKEAFREDL	407
Db	361	HYRNLHCALRPLDHESYEFKVISQYLQSTHAPTHSDYTMLLDLFEVEKDGEKEAFREDL	420
Qy	408	HNRMLLWHGSRMSNWWGILSHGLRIAPPEAPITGYMFGKGIYFADMSSKSANYCFASRLK	467
Db	421	HNRMLLWHGSRMSNWWGILSHGLRIAPPEAPITGYMFGKGIYFADMSSKSANYCFASRLK	480
Qy	468	NTGLLLLSEVALGQCNELLEANPKAEGLLQGKHSTKGLGKMAPSSAHFVTLNGSTVPLGP	527
Db	481	NTGLLLLSEVALGQCNELLEANPKAEGLLQGKHSTKGLGKMAPSSAHFVTLNGSTVPLGP	540
Qy	528	ASDTGILNPDGYTLNYNEYIVYNPNQVRMRYLLKVQFNFLQLW	570
Db	541	ASDTGILNPDGYTLNYNEYIVYNPNQVRMRYLLKVQFNFLQLW	583

RESULT 1

US-09-596-248D-2

; Sequence 2, Application US/09596248D

; Patent No. 6599727

; GENERAL INFORMATION:

; APPLICANT: Christenson, Erik

; APPLICANT: DeMaggio, Anthony J

; APPLICANT: Goldman, Phyllis S

; APPLICANT: McElligott, David L

; TITLE OF INVENTION: Human Poly(ADP-Ribose) Polymerase 2 Materials and

; TITLE OF INVENTION: Methods

; FILE REFERENCE: 27866/36544

; CURRENT APPLICATION NUMBER: US/09/596,248D

; CURRENT FILING DATE: 2000-06-16

; PRIOR APPLICATION NUMBER: 60/139,543

; PRIOR FILING DATE: 1999-06-16

; NUMBER OF SEQ ID NOS: 68

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 2

; LENGTH: 583

; TYPE: PRT

; ORGANISM: Homo sapiens

US-09-596-248D-2

Query Match 99.4%; Score 2981.5; DB 4; Length 583;
Best Local Similarity 97.8%; Pred. No. 2.8e-279;
Matches 570; Conservative 0; Mismatches 0; Indels 13; Gaps 1;

Qy	1	MAARRRRSTGGGRARALNESKRVNNGNTAPEDSSPAKKTTRCQRQESKKMPVAGGKANKD	60
Db	1	MAARRRRSTGGGRARALNESKRVNNGNTAPEDSSPAKKTTRCQRQESKKMPVAGGKANKD	60
Qy	61	RTEDKQD-----ESVKALLLKGPVDPECTAKVGKAHVYCEGNDVYDVMLN	107
Db	61	RTEDKQDGMPPGRSWASKRVSESVKALLLKGPVDPECTAKVGKAHVYCEGNDVYDVMLN	120
Qy	108	QTNLQFNNNKYYLIQLLEDDAQRNFSVWMRWGRVGKMGQHSLVACSGNLNKAKEIFQKKF	167
Db	121	QTNLQFNNNKYYLIQLLEDDAQRNFSVWMRWGRVGKMGQHSLVACSGNLNKAKEIFQKKF	180
Qy	168	LDKTKNNWEDREKFEKVPKGYDMLQMDYATNTQDEEETKKEESLKSPLKPESQLDLRVQE	227
Db	181	LDKTKNNWEDREKFEKVPKGYDMLQMDYATNTQDEEETKKEESLKSPLKPESQLDLRVQE	240
Qy	228	LIKLICNVQAMEEMMEMKYNTKKAPLGKLTVAQIKAGYQSLKKIEDCIRAGQHGRALME	287
Db	241	LIKLICNVQAMEEMMEMKYNTKKAPLGKLTVAQIKAGYQSLKKIEDCIRAGQHGRALME	300
Qy	288	ACNEFYTRIPHDFGLRTPPLIRTQKELSEKIQLLEALGDIEIAIKLVKTELQSPEHPLDQ	347
Db	301	ACNEFYTRIPHDFGLRTPPLIRTQKELSEKIQLLEALGDIEIAIKLVKTELQSPEHPLDQ	360
Qy	348	HYRNLHCALRPLDHESYEFKVISQYLQSTHAPTHSDYTMTLDDLFEVEKDGEKEAFREDL	407
Db	361	HYRNLHCALRPLDHESYEFKVISQYLQSTHAPTHSDYTMTLDDLFEVEKDGEKEAFREDL	420
Qy	408	HNRMLLWHGSRMSNWVGILSHGLRIAPPEAPITGYMFGKGIYFADMSSKSANYCFASRLK	467
Db	421	HNRMLLWHGSRMSNWVGILSHGLRIAPPEAPITGYMFGKGIYFADMSSKSANYCFASRLK	480
Qy	468	NTGLLLLSEVALGQCNELLEANPKAEGLLQGKHSTKGLGKMAPSSAHFVTLNGSTVPLGP	527
Db	481	NTGLLLLSEVALGQCNELLEANPKAEGLLQGKHSTKGLGKMAPSSAHFVTLNGSTVPLGP	540
Qy	528	ASDTGILNPDGYTLNNEYIVYNPNQVRMRYLLKVQFNFLQLW	570
Db	541	ASDTGILNPDGYTLNNEYIVYNPNQVRMRYLLKVQFNFLQLW	583